AMENDMENT TO THE CLAIMS

The following is a complete listing of the pending claims.

1. (Previously presented) A method for reducing the incorporation of norleucine into a

heterologous protein expressed by a microorganism comprising:

modifying a microorganism to co-express a heterologous protein and a non-standard

amino acid degrading protein, wherein the expression of the non-standard amino acid

degrading protein is increased relative to its expression in the microorganism before said

modifying step;

and wherein the non-standard amino acid degrading protein is a glutamate

dehydrogenase, leucine dehydrogenase, valine dehydrogenase, phenylalanine

dehydrogenase, or glutamate/leucine/phenylalanine/valine dehydrogenase.

2. (Previously presented) The method of claim 1 wherein the non-standard amino acid

degrading protein is a glutamate dehydrogenase.

3. (Currently amended) The method of claim 12 wherein the non-standard amino acid

degrading protein is a wild-type Escherichia coli glutamate dehydrogenase or a

Escherichia coli glutamate dehydrogenase having with a leucine at the amino acid

position that corresponds with amino acid position 92 of said wild-type glutamate

dehydrogenase, wherein the amino acid at position 92 of said wild-type glutamate

dehydrogenase is a lysine.

4. (Previously presented) The method of claim 3 wherein the non-standard amino acid

degrading protein comprises SEQ ID NO:2 or 4.

5. (Previously presented) The method of claim 4 wherein the non-standard amino acid

degrading protein is encoded by a DNA molecule comprising SEQ ID NO:1 or 3.

6-7. (Cancelled)

8. (Original) The method of claim 1 wherein the microorganism is *Escherichia coli*.

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Appl. No. 10/572,711 Response to Oct. 5, 2009 Final Office Action 9. (Previously presented) The method of claim 1 wherein the expressed heterologous

protein is a somatotropin.

10. (Previously presented) The method of claim 9 wherein the somatotropin is selected from

the group consisting of human, equine, bovine, ovine, porcine, canine, and feline

somatotropin.

- 11. (Original) The method of claim 9 wherein the somatotropin is bovine somatotropin.
- 12. (Previously presented) The method of claim 2 wherein the non-standard amino acid

degrading protein is an Escherichia coli glutamate dehydrogenase.

13. (Cancelled)

14. (Original) The method of claim 1 wherein the heterologous protein and the non-standard

amino acid degrading protein are expressed from a single expression vector.

15. (Original) The method of claim 1 wherein the heterologous protein and the non-standard

amino acid degrading protein are expressed from at least two distinct expression vectors.

16-41. (Cancelled)

42. (Previously presented) The method of claim 1 wherein the heterologous protein and/or

the non-standard amino acid degrading protein is expressed from a location in the

microorganism's genome.

43. (Previously presented) The method of claim 1 wherein the non-standard amino acid

degrading protein is a leucine dehydrogenase, a valine dehydrogenase, a

glutamate/leucine/phenylalanine/valine dehydrogenase, or a phenylalanine

dehydrogenase.

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- 44. (Withdrawn) The method of claim 43 wherein the non-standard amino acid degrading protein is a leucine dehydrogenase from *Bacillus cereus*, a leucine dehydrogenase from *Bacillus subtilis*, a leucine dehydrogenase from *Nostoc sp.*, a leucine dehydrogenase from *Shewanella oneidensis*, a valine dehydrogenase from *Streptomyces avermitilis*, or a glutamate/leucine/phenylalanine/valine dehydrogenase from *Nitrosomonas europaea*.
- 45. (Withdrawn) The method of claim 44 wherein the non-standard amino acid degrading protein comprises SEQ ID NO:6, 8, 10, 12, 14, or 16.
- 46. (Withdrawn) The method of claim 45 wherein the non-standard amino acid degrading protein is encoded by a DNA molecule comprising SEQ ID NO:5, 7, 9, 11, 13, or 15.

47-48. (Cancelled)

49. (Previously presented) The method of claim 1, wherein said non-standard amino acid degrading protein is a microbial non-standard amino acid degrading protein.